

AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions and listings of claims in the application.

Listing of Claims

Claim 1-56 (canceled).

Claim 57 (currently amended): A computer program ~~readable-medium-comprising computer-executable-instructions~~ for performing a method for acquiring data from a polymer array using an array scanner comprising:

~~scanning an area of a substrate having~~ a plurality of ~~different~~ diverse polymers of known sequence which form a polymer array that is located on a substrate,

wherein each ~~type of~~ diverse polymer has a different sequence and is on a localized area that is smaller than ~~250000~~ 2.5×10^5 microns²;

receiving pixel data of the polymer array from the scanner; and

collecting pixel data of the polymer array ; and

~~generating an average intensity for a given localized area.~~

Claim 58 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the method further comprises issuing commands to move a scan stage.

Claim 59 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the method further comprises outputting the pixel data to an image data file and displaying the image data.

Claim 60 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the polymers are nucleic acids and the substrate is hybridized with a sample.

Claim 61 (currently amended): The computer program ~~readable-medium~~ of Claim ~~60~~ 57 wherein the polymers are peptides.

Claim 62 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the substrate has at least 400 polymers per cm².

Claim 63 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the substrate has at least 1000 polymers per cm².

Claim 64 (currently amended): The computer program ~~readable-medium~~ of Claim 57 wherein the substrate has at least 10000 polymers per cm².

Claim 65 (currently amended): A ~~computer~~ Computer software ~~product~~ comprising:
 ~~computer program~~ software code that scans ~~an area of a substrate having~~ a plurality of ~~different~~ diverse polymers of known sequence which form an array that is located on a substrate, wherein each ~~type of~~ diverse polymer has a different sequence and is on a localized area ~~areas~~ that is smaller than ~~250000~~ 2.5 x 10⁵ microns²;
 ~~computer program~~ software code that receives pixel data of the polymer array from the scanner;
 ~~computer program~~ software code that collects pixel data of the polymer array ;
 ~~computer program code that generates an average intensity for a given localized area;~~
and
 ~~a computer readable medium for storing the codes .~~

Claim 66 (currently amended): The computer software ~~product~~ of Claim 65 further comprising ~~computer program~~ software code that issues commands to move a scan stage.

Claim 67 (currently amended): The computer software ~~product~~ of Claim 66 further comprising ~~computer program~~ software code that outputs the pixel data to an image data file and displays the image data.

Claim 68 (currently amended): The computer software ~~product~~ of Claim 65 wherein the polymers are nucleic acids and the substrate is hybridized with a sample.

Claim 69 (currently amended): The computer software ~~product~~ of Claim 65 wherein the polymers are peptides.

Claim 70 (currently amended): The computer software ~~product~~ of Claim 65 wherein the substrate has at least 400 polymers per cm².

Claim 71 (currently amended): The computer software ~~product~~ of Claim 65 wherein the substrate has at least 1000 polymers per cm².

Claim 72 (currently amended): The computer software ~~product~~ of Claim 65 wherein the substrate has at least 10000 polymers per cm².

Claim 73 (currently amended): A system for acquiring data from an array of diverse polymer sequences using a polymer scanner comprising:

~~a processor; and a memory being coupled to the processor, the memory storing a plurality of machine instructions that cause the processor to perform~~ a computer containing a program that implements a plurality of ~~logical~~ steps ~~when implemented by the processor~~ , the ~~logical~~ steps comprising:

~~scanning an area of a substrate having a plurality of different~~ diverse polymers of known sequence which form an array that is located on a region of a substrate, wherein each ~~type of diverse polymer has a different sequence and~~ is on a localized area that is smaller than ~~250000~~ 2.5 x 10⁵ microns²;

receiving pixel data of the polymer array from the scanner; and

collecting pixel data of the polymer array ;~~and~~

~~generating an average intensity for a given localized area.~~

Claim 74 (currently amended): The system of Claim 73 wherein the ~~logic~~ steps further ~~comprises~~ comprise issuing commands to move a scan stage.

Claim 75 (currently amended): The system of Claim 74 wherein the ~~logic~~ steps further ~~comprises~~ comprise outputting the pixel data to an image data file and displaying the image data.

Claim 76 (previously presented): The system of Claim 75 wherein the polymers are nucleic acids and the substrate is hybridized with a sample.

Claim 77 (previously presented): The system of Claim 76 wherein the polymers are peptides.

Claim 78 (currently amended): A computer program ~~readable-medium-comprising computer-executable-instructions~~ for performing a method comprising:

scanning a plurality of diverse polymer probes of known sequence which form a polymer array to obtain a plurality of intensity data, wherein the polymer array has a plurality of ~~different~~ diverse polymer probes of known sequence on a substrate, wherein each diverse type of polymer probe has a different sequence and occupies a localized area that is less than 250000 2.5 x 10⁵ microns², and wherein the array has been contacted with a sample that may contain a target; and

determining the positions of probe and target interaction based upon the intensity data of the polymer array; and

collecting pixel data of the polymer array and

~~generating an average intensity for a given localized area.~~

Claim 79 (currently amended): The computer program ~~readable-medium~~ of Claim 78 wherein the polymer probes are nucleic acids.

Claim 80 (currently amended): The computer program ~~readable-medium~~ of Claim 78 wherein the polymer probes are oligonucleotides.

Claim 81 (currently amended): The computer program ~~readable-medium~~ of Claim 80 wherein the intensity data reflects the hybridization of the oligonucleotide probes and the target.

Claim 82 (currently amended): The computer program ~~readable-medium~~ of Claim 78 further comprising generating an average intensity based on ~~wherein intensity data are~~ fluorescence ~~fluorescence~~ data.

Claim 83 (currently amended): The computer program ~~readable-medium~~ of Claim 78 wherein the substrate has at least 400 polymer probes per cm².

Claim 84 (currently amended): The computer program ~~readable-medium~~ of Claim 78 wherein the substrate has at least 1000 polymer probes per cm².

Claim 85 (currently amended): The computer program ~~readable-medium~~ of Claim 78 wherein the substrate has at least 10000 polymer probes per cm².

Claim 86 (currently amended): Computer A-computer software product comprising:

~~computer program~~ software code that scans a ~~polymer array to obtain a plurality~~ intensity data wherein the polymer array has a plurality of different diverse polymer probes, of known sequence which form an array on a substrate, wherein each ~~type of~~ diverse polymer probe has a different sequence and occupies a localized area that is less than ~~250000~~ 2.5 x 10⁵ microns², and wherein the array has been contacted with a sample that may contain a target; and

~~computer program~~ software code that determines the positions of probe and target interaction ~~based upon the intensity data and the computer program code collects pixel data and generates an average intensity for a given localized area ; and~~

~~a computer readable medium for storing the codes .~~

Claim 87 (currently amended): The computer software ~~product~~ of Claim 86 wherein the polymer probes are nucleic acids.

Claim 88 (currently amended): The computer software ~~product~~ of Claim 86 wherein the polymer probes are oligonucleotides.

Claim 89 (currently amended): The computer software ~~product~~ of Claim 88 wherein the intensity data reflects the hybridization of the oligonucleotide probes and the target.

Claim 90 (currently amended): The computer software ~~product~~ of Claim 86 wherein intensity data are ~~floreseence~~ fluorescence data.

Claim 91 (currently amended): The computer software ~~product~~ of Claim 86 wherein the substrate has at least 400 polymer probes per cm^2 .

Claim 92 (currently amended): The computer software ~~product~~ of Claim 86 wherein the substrate has at least 1000 polymer probes per cm^2 .

Claim 93 (currently amended): The computer software ~~product~~ of Claim 86 wherein the substrate has at least 10000 polymer probes per cm^2 .

Claim 94 (currently amended): A system for acquiring data from an array having diverse polymer sequences using a polymer scanner comprising:

~~a processor; and a memory being coupled to the processor, the memory storing a plurality of machine instructions that cause the processor to perform~~ a computer containing a program that implements a plurality of logical steps ~~when implemented by the processor, the logical steps comprising:~~

scanning a ~~polymer array to obtain a plurality of intensity data, wherein the polymer array has~~ a plurality of ~~different~~ diverse polymer probes of known sequence which form an array on a substrate, wherein each ~~type of~~ diverse polymer probe has a different sequence and occupies a localized area that is less than 250000 2.5×10^5 microns², and wherein the array has been contacted with a sample that may contain a target; and

determining the positions of probe and target interaction based upon fluorescent intensity data of the polymer array; and

collecting pixel data and

~~generating an average intensity for a given localized area.~~

Claim 95 (previously presented): The system of Claim 94 wherein the polymer probes are nucleic acids.

Claim 96 (previously presented): The system of Claim 95 wherein the polymer probes are oligonucleotides.

Claim 97 (previously presented): The system of Claim 96 wherein the intensity data reflects the hybridization of the oligonucleotide probes and the target.

Claim 98 (currently amended): The system of Claim 97 further comprising generating an average intensity based on ~~wherein intensity data are florescence~~ fluorescence data.

Claim 99 (previously presented): The system of Claim 94 wherein the substrate has at least 400 polymer probes per cm².

Claim 100 (previously presented): The system of Claim 94 wherein the substrate has at least 1000 polymer probes per cm².

Claim 101 (previously presented): The system of Claim 94 wherein the substrate has at least 10000 polymer probes per cm².

Claim 102 (currently amended): A system for scanning a polymer array comprising:

a scanning optical device;

a polymer array having ~~different~~ diverse polymers of known sequence on a substrate wherein each ~~type of diverse~~ polymer has a different sequence and is in a localized area that is smaller than 2.5×10^5 microns²;

~~a processor; and a memory being coupled to the processor, the memory storing a plurality of machine instructions that cause the processor to perform~~ a computer containing a program that implements a plurality of ~~logical steps when implemented by the processor, the logical step~~ steps comprising collecting fluorescent intensity data from less than 1/2 of each of the localized areas from the polymer array and generating an average intensity for at least one of said a given localized areas area from the polymer array.

Claim 103 (previously presented): The system of Claim 102 wherein the polymer array is a nucleic acid probe array.

Claim 104 (previously presented): The system of Claim 103 wherein the polymer array is a peptide array.

Claim 105 (previously presented): The system of Claim 104 wherein the polymer array has at least 400 polymers per cm².

Claim 106 (previously presented): The system of Claim 105 wherein the polymer array has at least 1000 polymers per cm².

Claim 107 (previously presented): The system of Claim 106 wherein the polymer array has at least 10000 polymers per cm^2 .

Claim 108 (currently amended): A computer program ~~readable-medium comprising executable instructions~~ for acquiring data from a polymer array, comprising:

scanning ~~a substrate having~~ a plurality of ~~different~~ diverse polymers, of known sequence which from an array on a substrate, wherein each ~~type of~~ diverse polymer is has a different sequence and is on in a localized area, ~~having an area that is~~ smaller than ~~250000~~ 2.5 $\times 10^5$ microns²,

acquiring data from the polymer array which indicate binding between the polymer on the substrate and a detectable target polymer; and

collecting pixel data of the polymer array; ~~and~~

~~generating an average intensity for a given localized area.~~

Claim 109 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the target polymer is a polypeptide.

Claim 110 (currently amended): The computer program ~~readable-medium~~ of Claim 109 wherein the ~~solid~~ substrate has at least 400 probe polymers per cm^2 .

Claim 111 (currently amended): The computer program ~~readable-medium~~ of Claim 110 wherein the ~~solid~~ substrate has at least 1000 probe polymers per cm^2 .

Claim 112 (currently amended): The computer program ~~readable-medium~~ of Claim 111 wherein the ~~solid~~ substrate has at least 10,000 ~~probe~~ polymers per cm^2 .

Claim 113 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the target polymer is a nucleic acid.

Claim 114 (currently amended): The computer program ~~readable-medium~~ of Claim 113 wherein the ~~solid~~ substrate has at least 400 ~~probe~~ polymers per cm^2 .

Claim 115 (currently amended): The computer program ~~readable-medium~~ of Claim 114 wherein the ~~solid~~ substrate has at least 1000 ~~probe~~ polymers per cm^2 .

Claim 116 (currently amended): The computer program ~~readable-medium~~ of Claim 115 wherein the ~~solid~~ substrate has at least 10,000 ~~probe~~ polymers per cm^2 .

Claim 117 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the data are fluorescence intensities.

Claim 118 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the substrate is an impermeable substrate having at least 1000 polymers/ cm^2 .

Claim 119 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein each of the localized areas ~~known-locations~~ is smaller than 1×10^4 ~~10,000 or 2,500~~ microns².

Claim 120 (currently amended): A computer software ~~product~~ comprising:

computer-program software code that scans a-substrate-having a plurality of different diverse nucleic acids or polypeptides polymers of known sequence which form an array on a substrate, wherein each diverse nucleic acid or polypeptide has a different sequence and is on

a at localized area areas, ~~each of which~~ that is an area smaller than ~~250000~~ 2.5×10^5 microns²; and

~~computer program~~ software code that acquires data from the array which indicate binding between the nucleic acid or polypeptide polymer on the substrate and a detectable target ~~polymer~~ and collects pixel data of the array ~~and generates an average intensity for individual localized areas; and~~

~~a computer readable medium for storing the codes .~~

Claim 121 (canceled)

Claim 122 (currently amended): The computer software ~~product~~ of Claim 121 wherein the ~~solid~~ substrate has at least 400 ~~probe polymers~~ nucleic acids or polypeptides per cm².

Claim 123 (currently amended): The computer software ~~product~~ of Claim 122 wherein the ~~solid~~ substrate has at least 1000 ~~probe polymers~~ nucleic acids or polypeptides per cm².

Claim 124 (currently amended): The computer software ~~product~~ of Claim 123 wherein the ~~solid~~ substrate has at least 10,000 ~~probe polymers~~ nucleic acids or polypeptides per cm².

Claim 125 (currently amended): The computer software ~~product~~ of Claim 120 wherein the target ~~polymer~~ is a nucleic acid.

Claim 126 (canceled)

Claim 127 (canceled)

Claim 128 (canceled)

Claim 129 (currently amended): The computer software ~~product~~ of Claim 120 wherein the data are fluorescence intensities.

Claim 130 (currently amended): The computer software ~~product~~ of Claim 120 wherein the substrate is an impermeable substrate having at least 1000 nucleic acids or polypeptides polymers/cm².

Claim 131 (currently amended): The computer software ~~product~~ of Claim 120 wherein each of the ~~known locations~~ localized areas is smaller than 1 x 10⁴ ~~10,000 or 2,500~~ microns².

Claim 132 (currently amended): A computer program ~~readable medium comprising executable instructions~~ for acquiring data from a ~~polymer~~ nucleic acid or polypeptide array, comprising:

~~scanning a substrate having~~ a plurality of ~~different~~ diverse polymers nucleic acids or polypeptides of known or detectable sequence ~~at localized areas~~ which form an array on a substrate, each diverse nucleic acid or polypeptide having a different sequence and is on each of which a localized area that is are smaller than ~~250000~~ 2.5 x 10⁵ microns²; and

acquiring data from the array which indicate binding between the nucleic acids or polypeptides ~~polymer~~ on the substrate and a detectable target ~~polymer~~; and

collecting pixel data of the array; ~~and~~

~~generating an average intensity for a given localized area.~~

Claim 133 (canceled)

Claim 134 (currently amended): The computer program ~~readable-medium~~ of Claim 109 wherein the ~~solid~~ substrate has at least 400 ~~probe polymers~~ nucleic acids or polypeptides per cm^2 .

Claim 135 (currently amended): The computer program ~~readable-medium~~ of Claim 110 wherein the ~~solid~~ substrate has at least 1000 ~~probe polymers~~ nucleic acids or polypeptides per cm^2 .

Claim 136 (currently amended): The computer program ~~readable-medium~~ of Claim 111 wherein the ~~solid~~ substrate has at least 10,000 ~~probe polymers~~ nucleic acids or polypeptides per cm^2 .

Claim 137 (canceled)

Claim 138 (canceled)

Claim 139 (canceled)

Claim 140 (canceled)

Claim 141 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the data are fluorescence intensities.

Claim 142 (currently amended): The computer program ~~readable-medium~~ of Claim 108 wherein the substrate is an impermeable substrate having at least 1000 nucleic acids or polypeptides polymers/ cm^2 .

Claim 143 (currently amended): The computer program ~~readable medium~~ of Claim 108 wherein each of the ~~known locations~~ localized areas is smaller than 1×10^4 ~~10,000 or 2,500~~ microns².

Claim 144 (currently amended): A system for controlling a ~~polymer~~ scanner for a nucleic acid array comprising:

~~a processor; and a memory being coupled to the processor, the memory storing a plurality of machine instructions that cause the processor to perform~~ a computer containing a program that implements a plurality of logical steps ~~when implemented by the processor, the logical steps comprising:~~

scanning ~~a substrate having~~ a plurality of ~~different~~ diverse nucleic acid polymers; of known or detectable sequence, ~~at localized areas on a substrate, each diverse nucleic acid polymer having a different sequence and on each of which are~~ a localized area that is smaller than ~~250000~~ 1×10^4 microns²; and

acquiring fluorescent data from the nucleic acid array which indicate binding between the ~~polymer~~ nucleic acid on the substrate and a detectable target ~~polymer~~;

collecting pixel data of the nucleic acid array; and

~~generating an average intensity for a localized area.~~

Claim 145 (canceled)

Claim 146 (canceled)

Claim 147 (canceled)

Claim 148 (canceled)

Claim 149 (previously presented): The system of Claim 144 wherein the target ~~polymer~~ is a nucleic acid.

Claim 150 (previously presented): The system of Claim ~~149~~ 144 wherein the ~~solid~~ substrate has at least 400 ~~probe polymers~~ nucleic acids per cm².

Claim 151 (previously presented): The system of Claim ~~150~~ 144 wherein the solid substrate has at least 1000 ~~probe polymers~~ nucleic acids per cm².

Claim 152 (previously presented): The system of Claim ~~151~~ 144 wherein the solid substrate has at least 10,000 ~~probe polymers~~ nucleic acids per cm².

Claim 153 (canceled)

Claim 154 (previously presented): The system of Claim 144 wherein the substrate is an impermeable substrate having at least 1000 ~~polymers~~ nucleic acids/cm².

Claim 155 (canceled)

Claim 156 (new): The computer program of claim 60 wherein the pixel data is from fluorescent intensity data.

Claim 157 (new): The computer software of claim 68 wherein the pixel data is from fluorescent intensity data.

Claim 158 (new): The system of claim 76 wherein the pixel data is from fluorescent intensity data.

Claim 159 (new): The computer program of claim 79 wherein the intensity data is fluorescent intensity.

Claim 160 (new): The computer software of claim 87 wherein the pixel data is from fluorescent intensity data.

Claim 161 (new): The computer software of claim 120 wherein the pixel data is from fluorescent intensity data.

Claim 162 (new): The computer program of claim 132 wherein the pixel data is from fluorescent intensity data.